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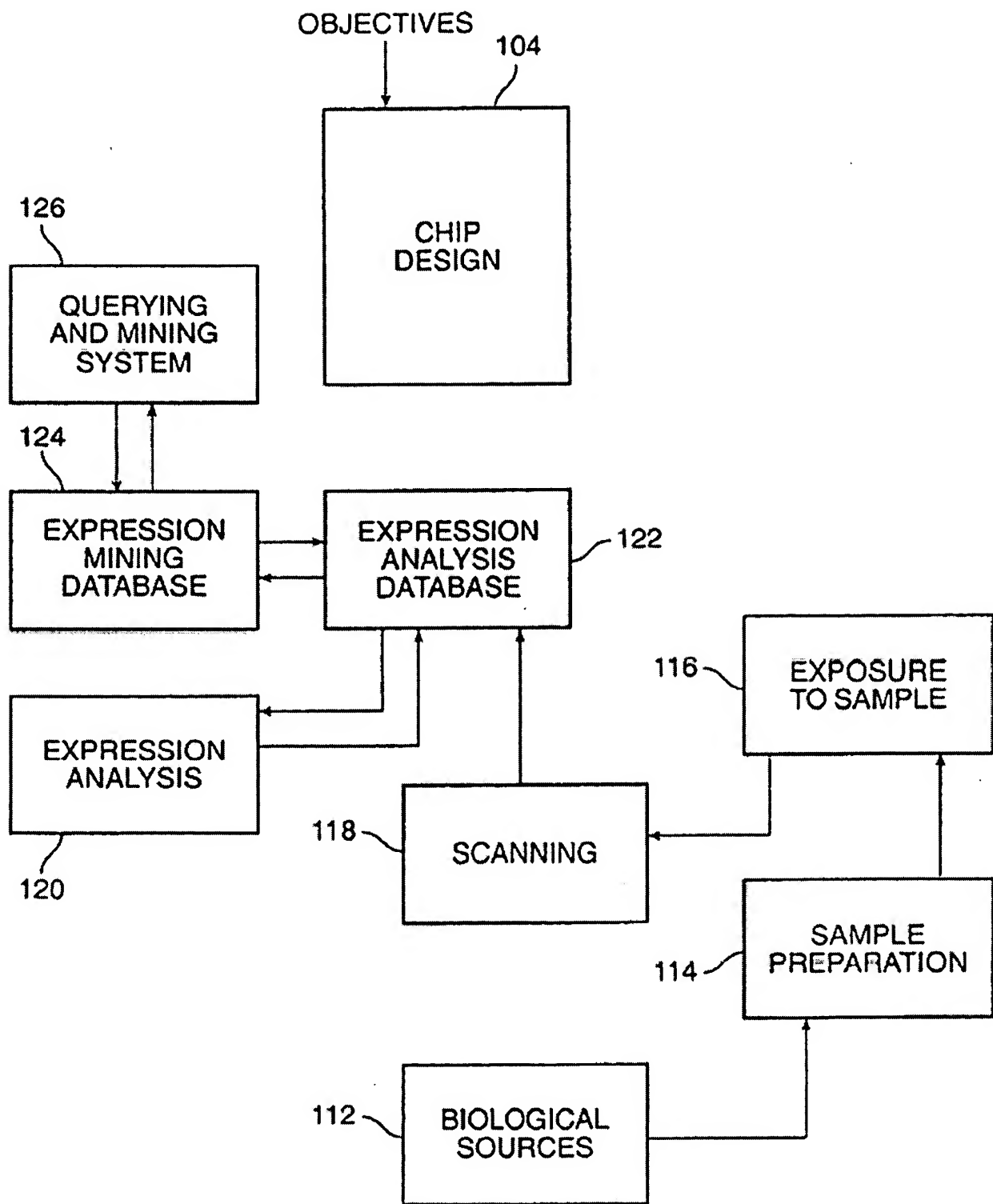


FIG. 1

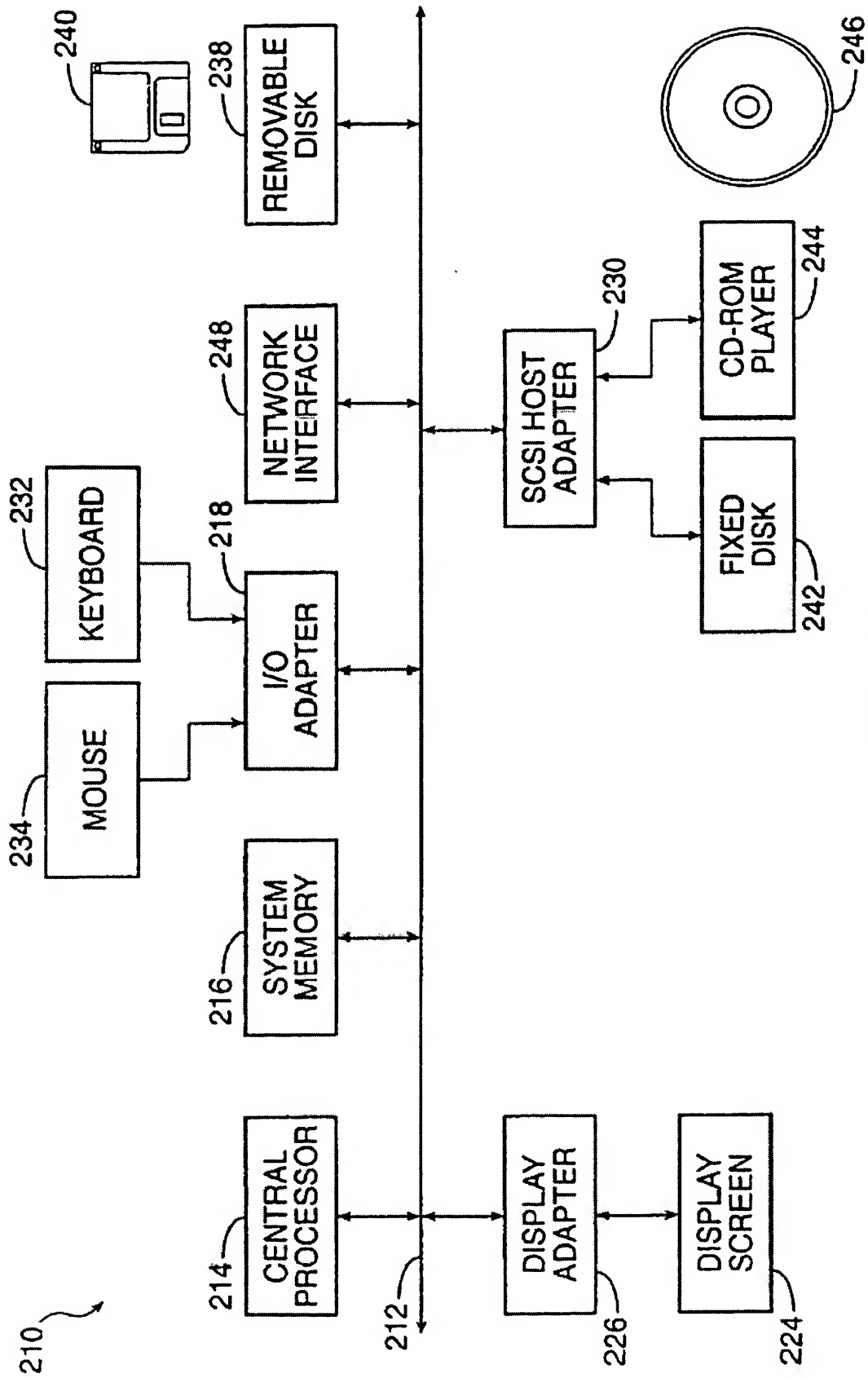


FIG. 2A

260 ~~~~~

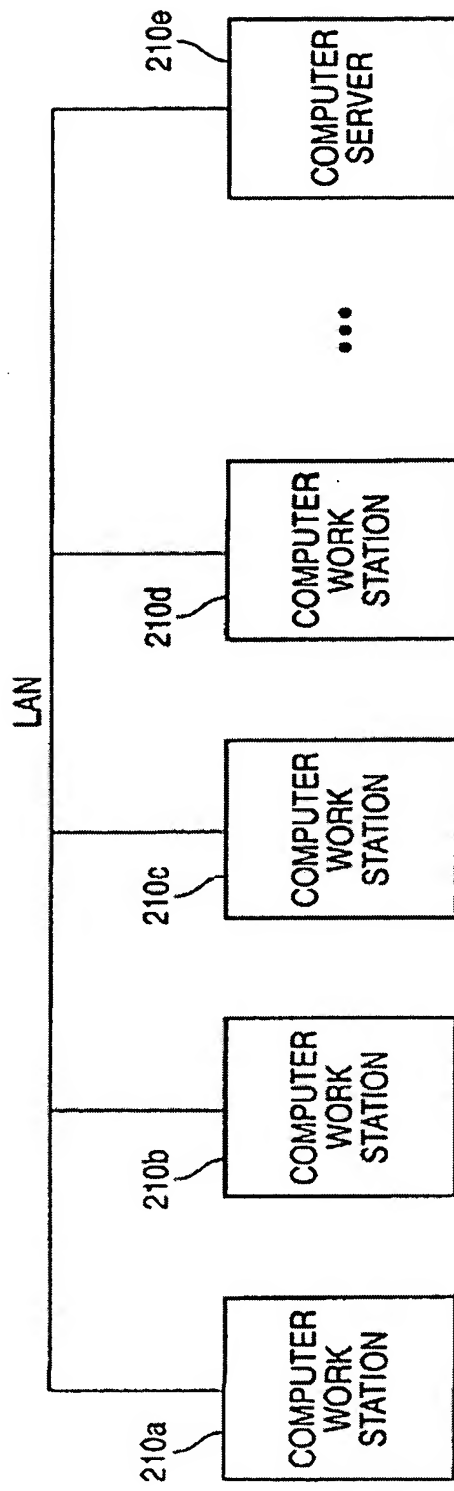


FIG. 2B

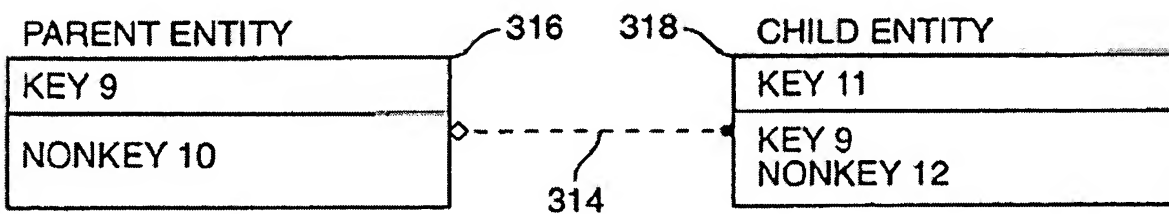
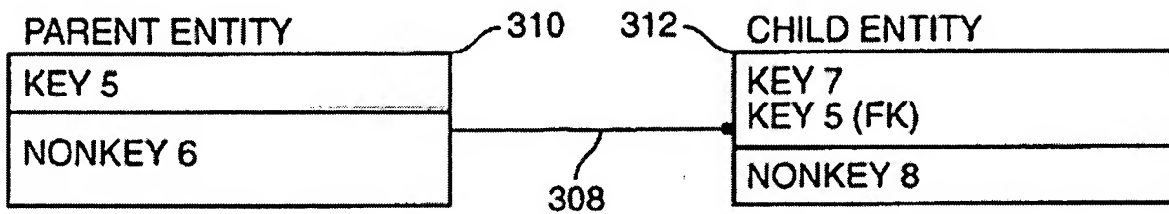
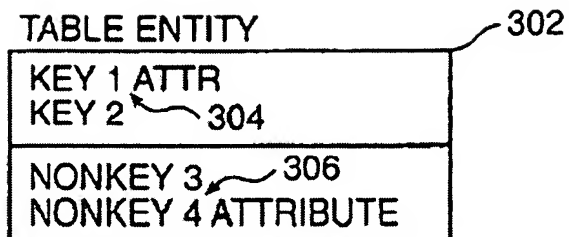


FIG. 3

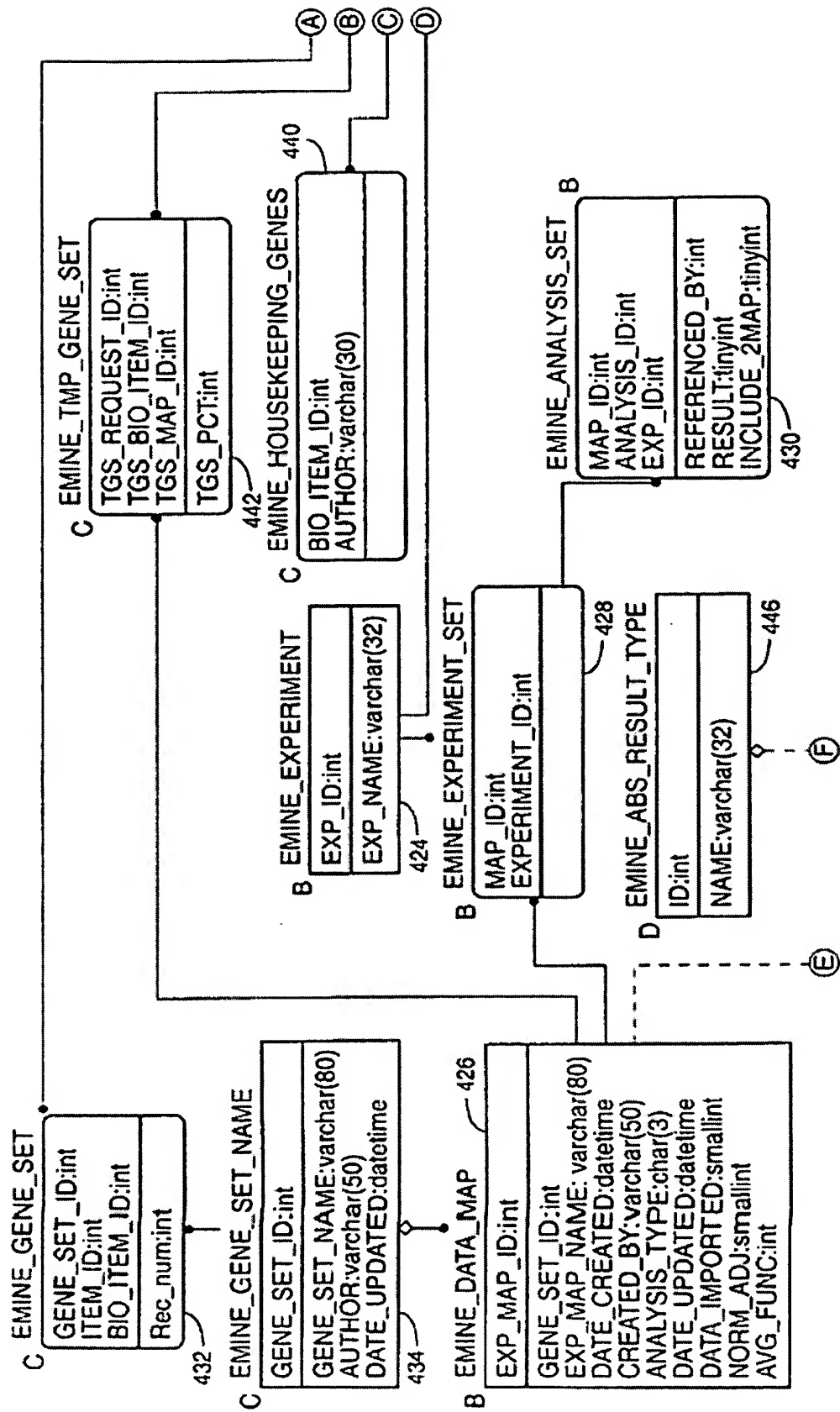


FIG. 4A.1

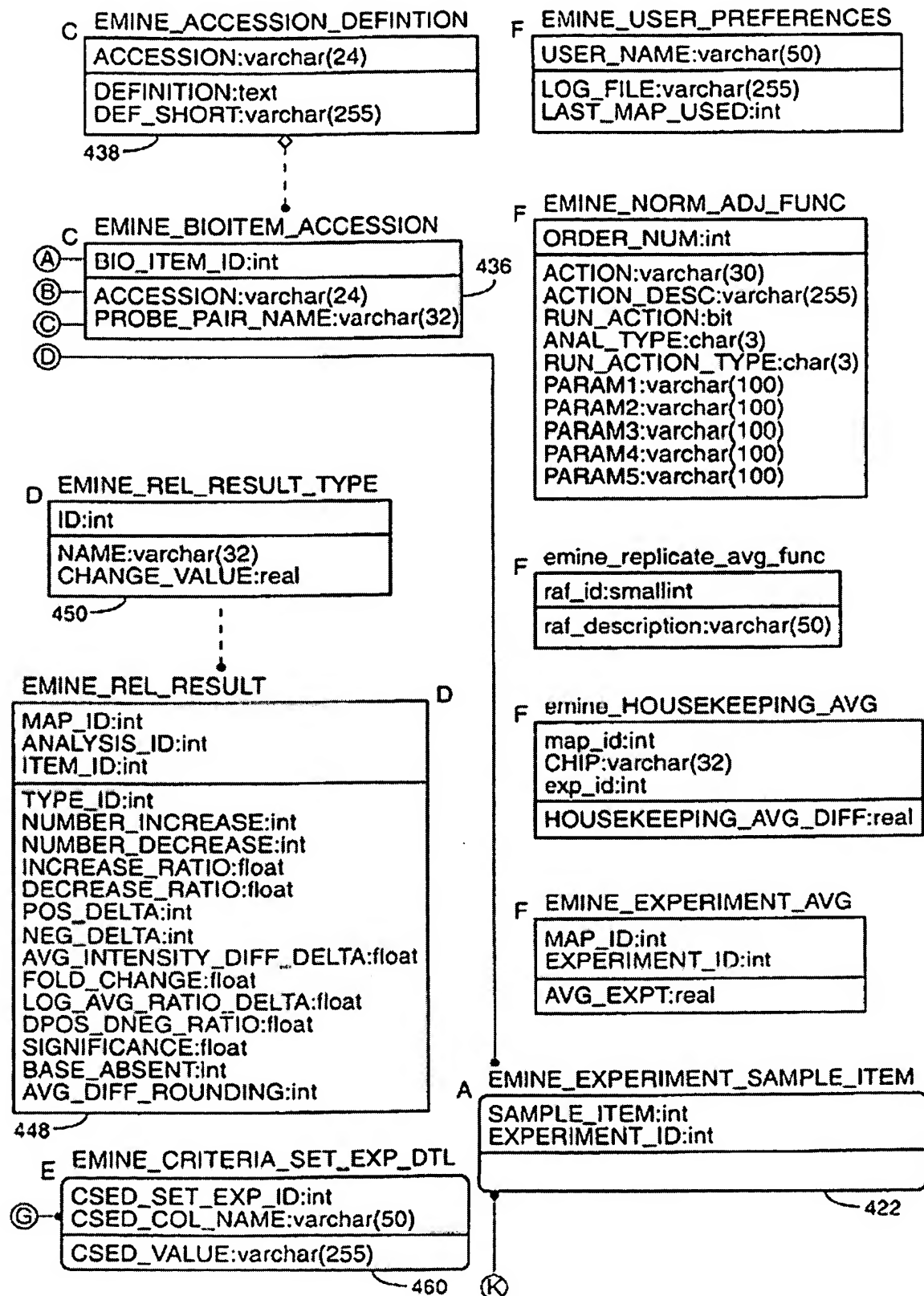


FIG. 4A.3

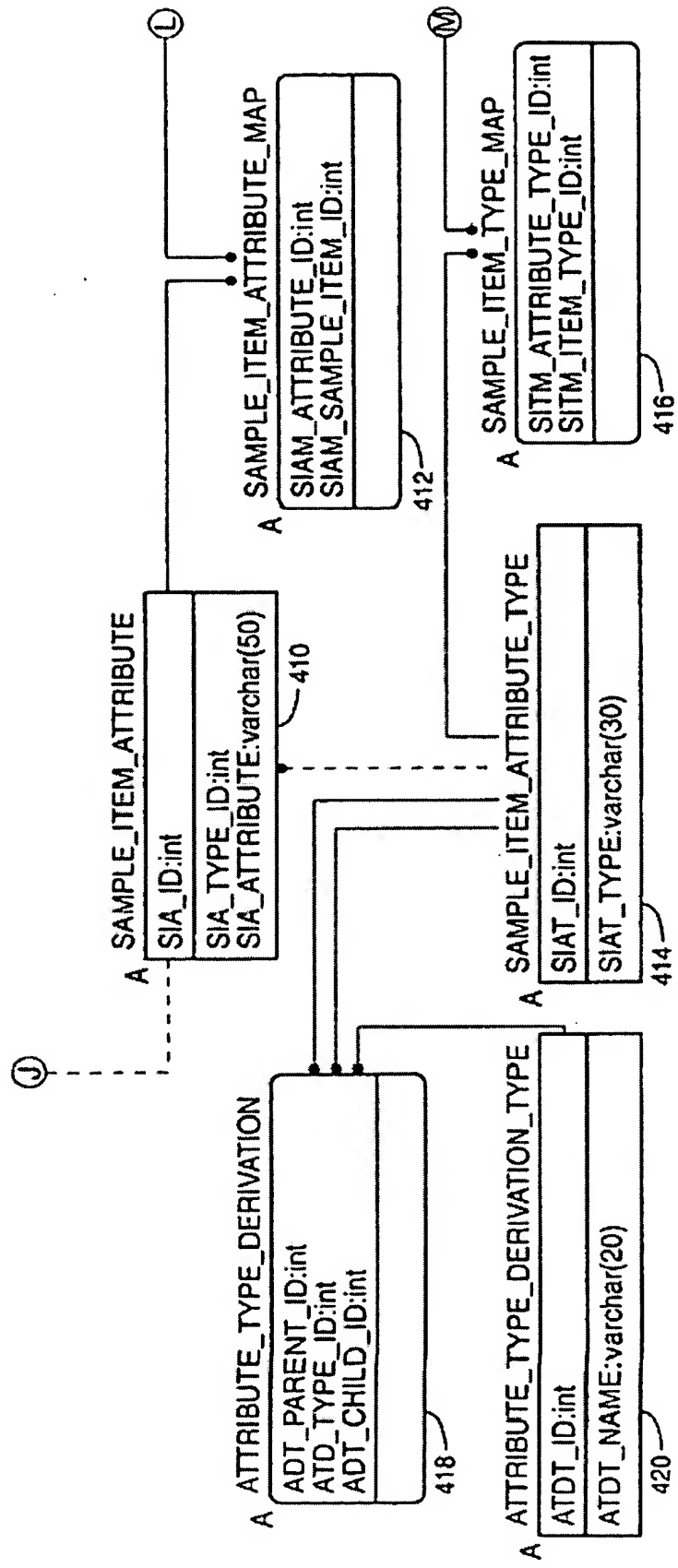


FIG. 4A.4

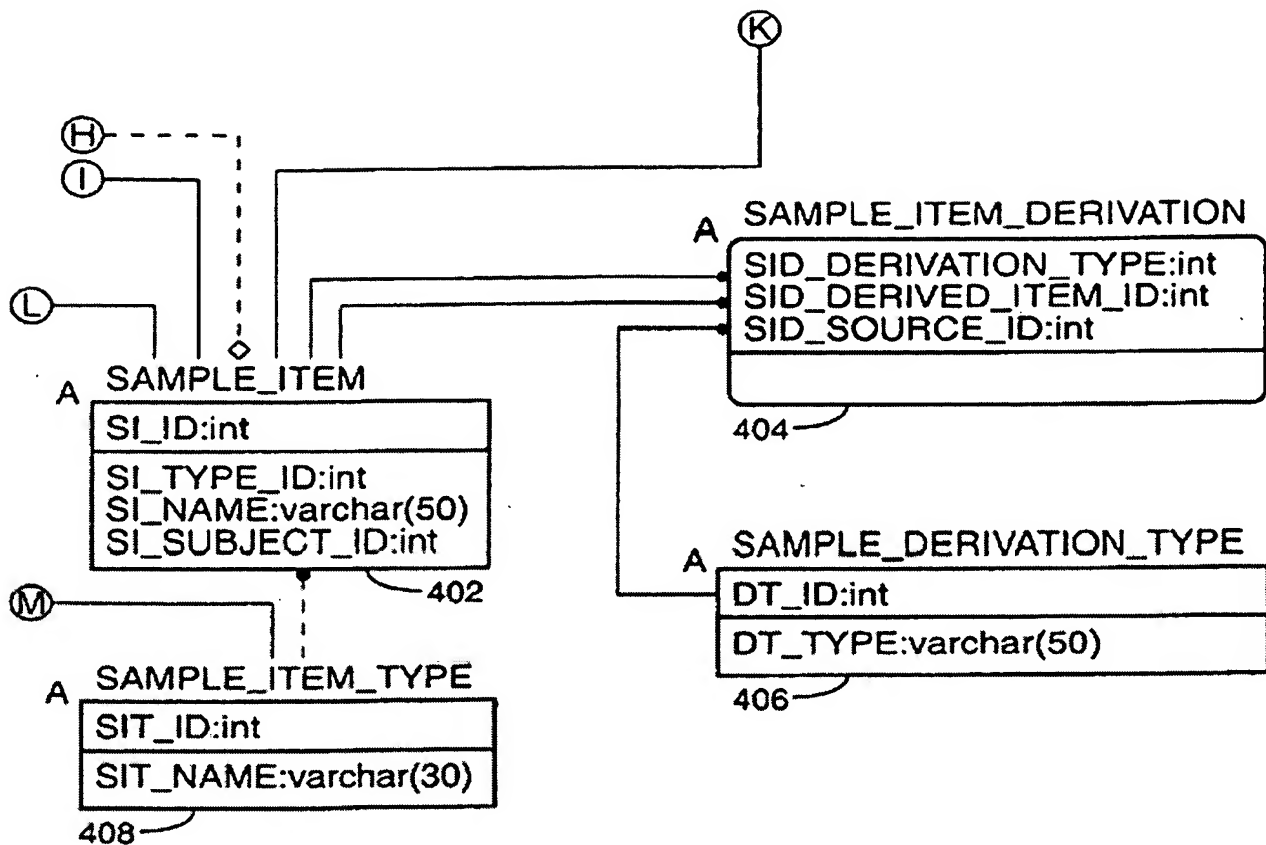


FIG. 4A.5

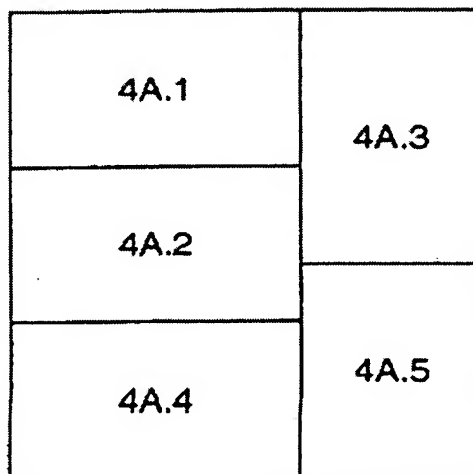


FIG. 4A

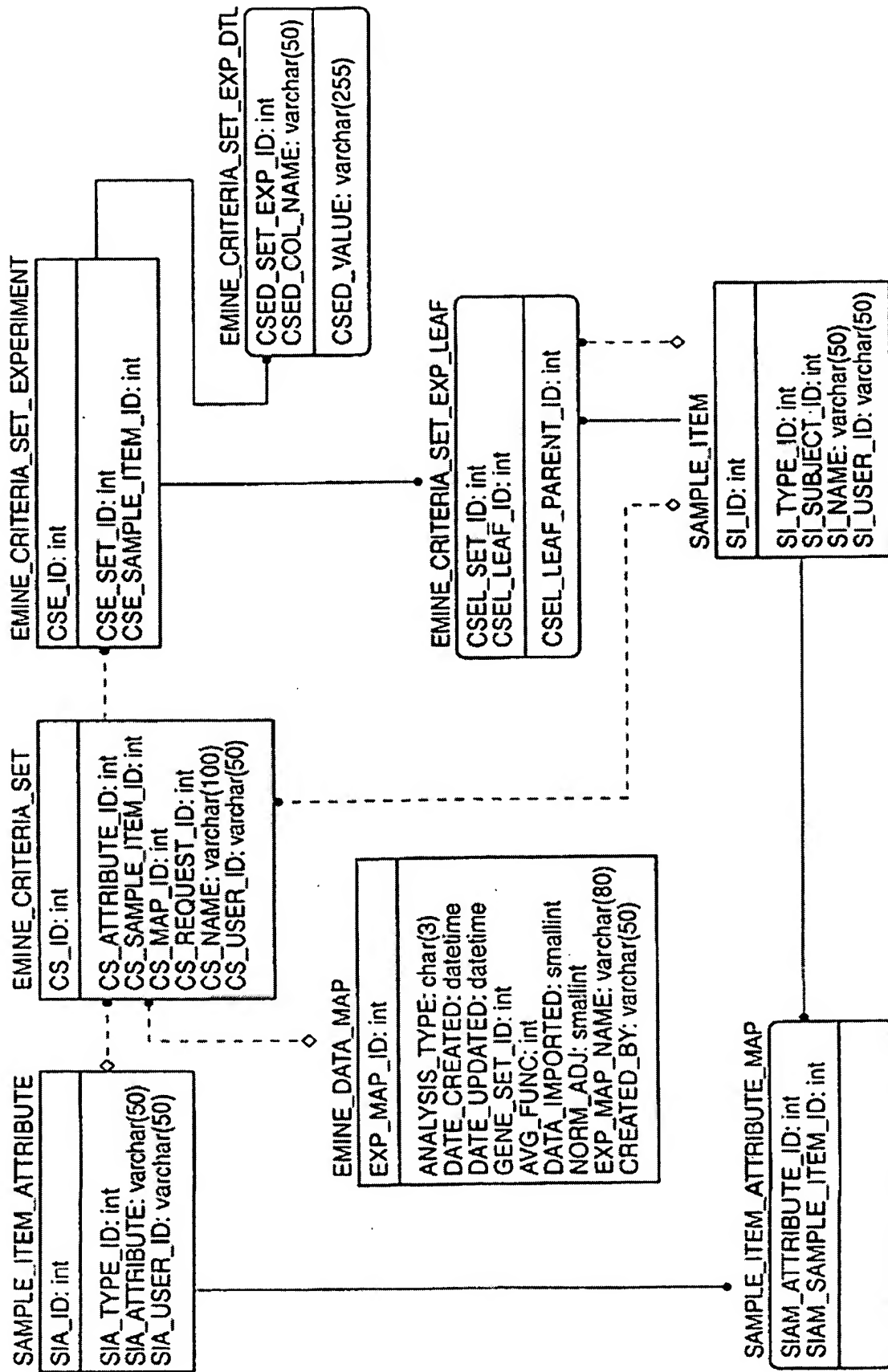


FIG. 4B

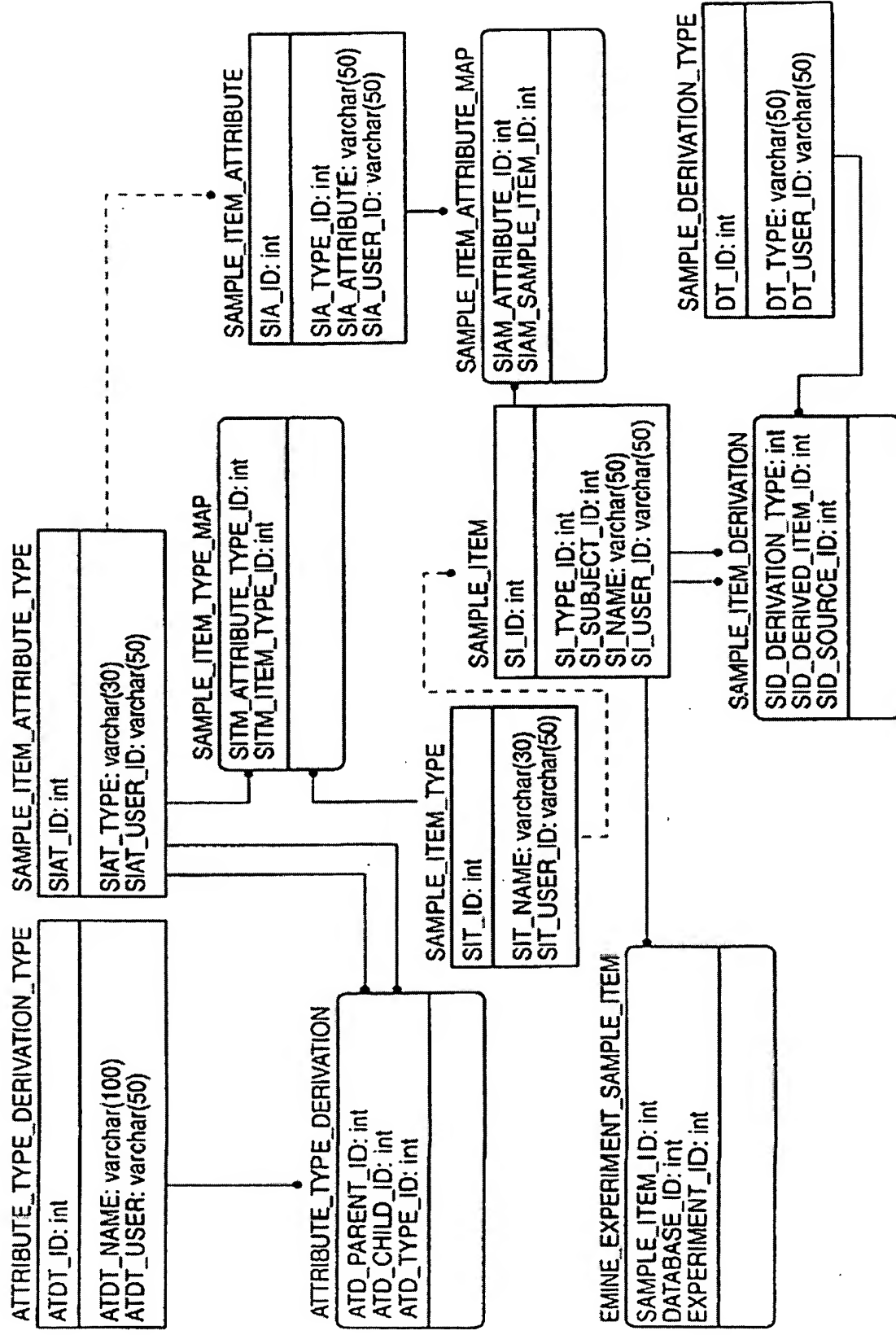


FIG. 4C

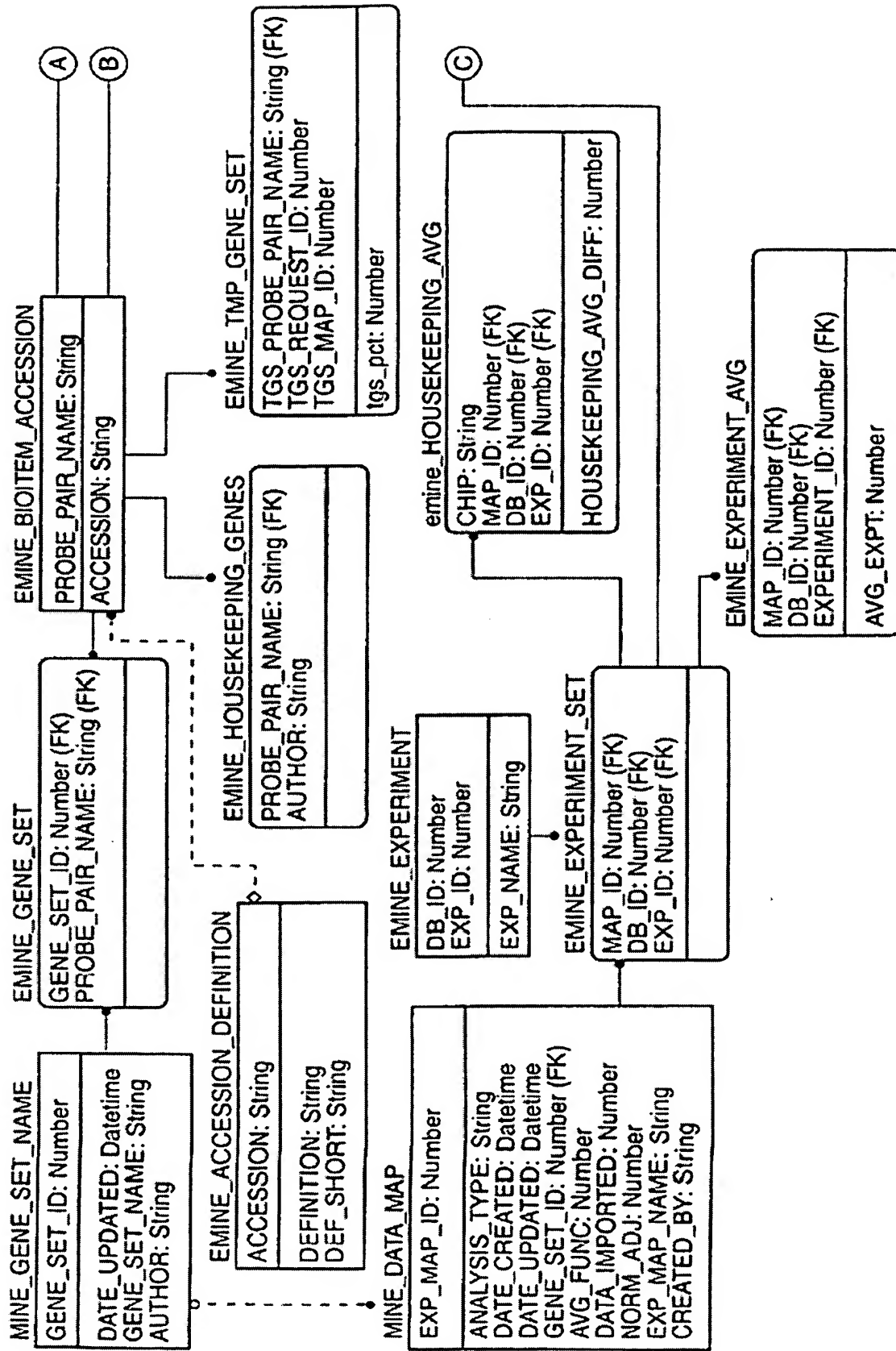


FIG. 4D.1

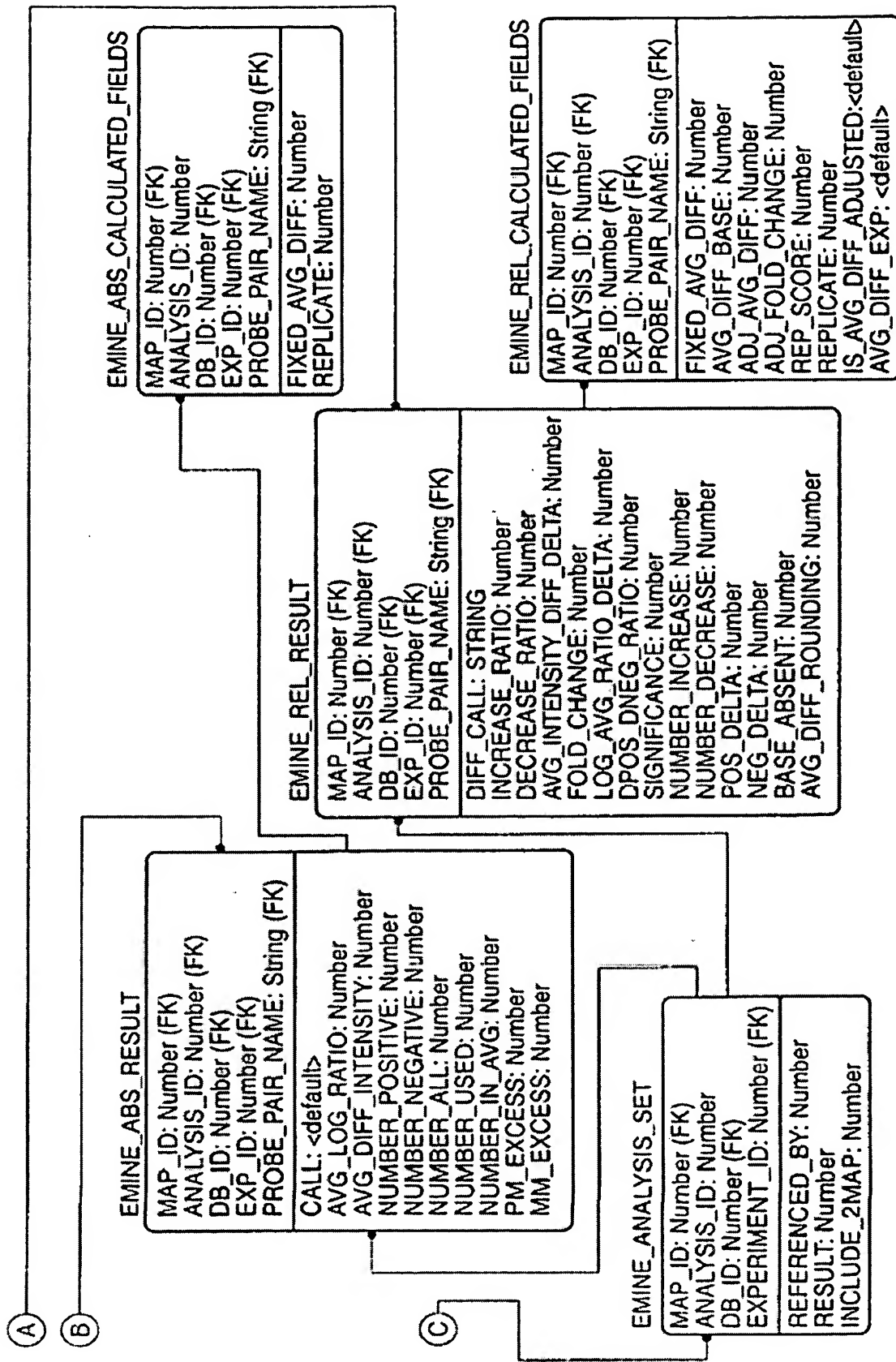


FIG. 4D.2

| EMINE_REPLICATE_AVG_FUNC | |
|--------------------------|--|
| RAF_ID | |
| RAF_DESCRIPTION | |

| EMINE_NORM_ADJ_FUNC | |
|---------------------|--|
| ORDER_NUM | |
| RUN_ACTION | |
| ANAL_TYPE | |
| RUN_ACTION_TYPE | |
| ACTION | |
| ACTION_DESC | |
| PARAM1 | |
| PARAM2 | |
| PARAM3 | |
| PARAM4 | |
| PARAM5 | |

| EMINE_USER_PREFERENCES | |
|------------------------|--|
| USER_NAME | |
| UPDATE_BIO_REF | |
| LAST_MAP_USED | |
| LAST_DB_ID | |
| LOG_FILE | |

| EMINE_SELECTED_SAMPLE_ITEMS | |
|-----------------------------|--|
| SSI_MAP_ID | |
| SSI_SAMPLE_ITEM_ID | |
| | |

| EMINE_TMP_SAMPLE_ITEM_ID | |
|--------------------------|--|
| TSI_MAP_ID | |
| TSI_SI_ID | |
| | |

FIG. 4E

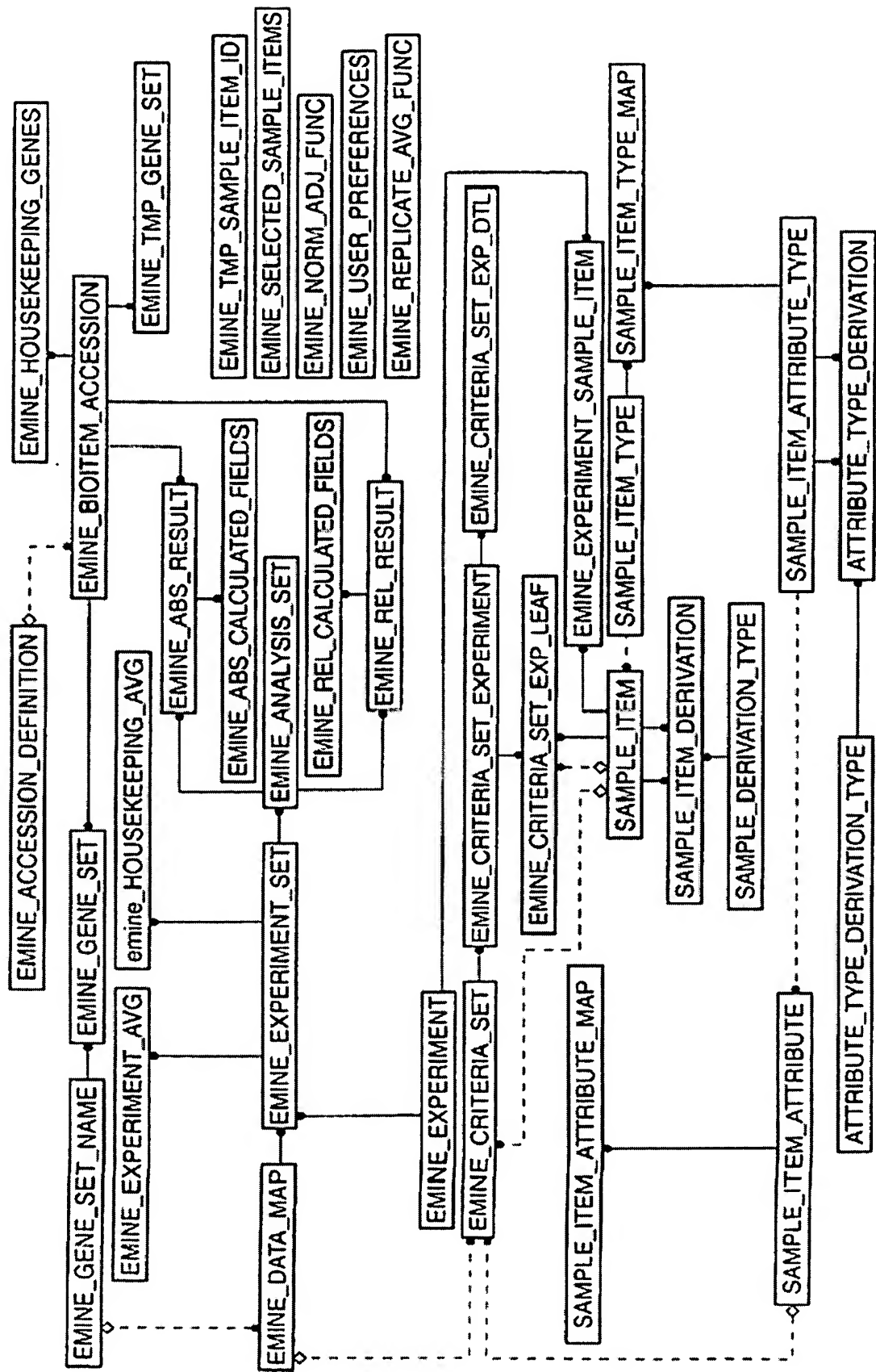


FIG. 4F

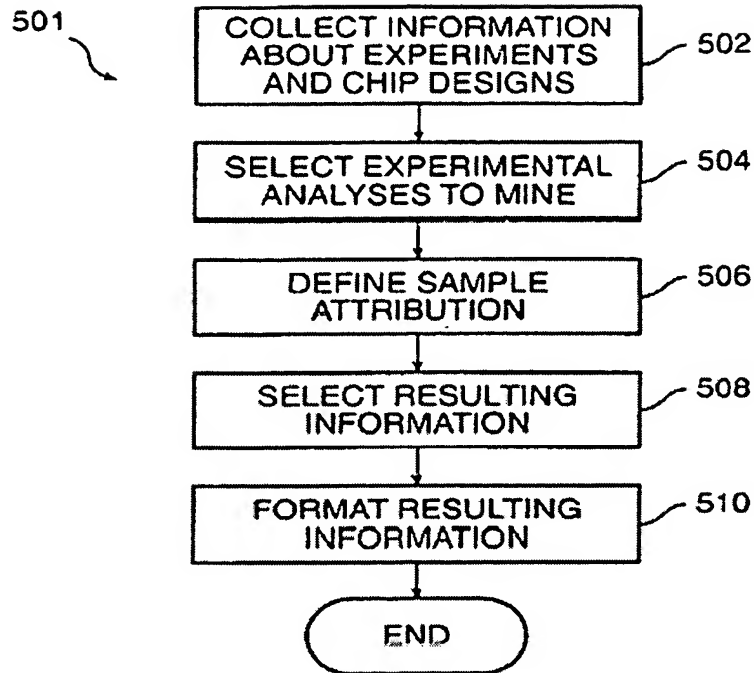


FIG. 5A

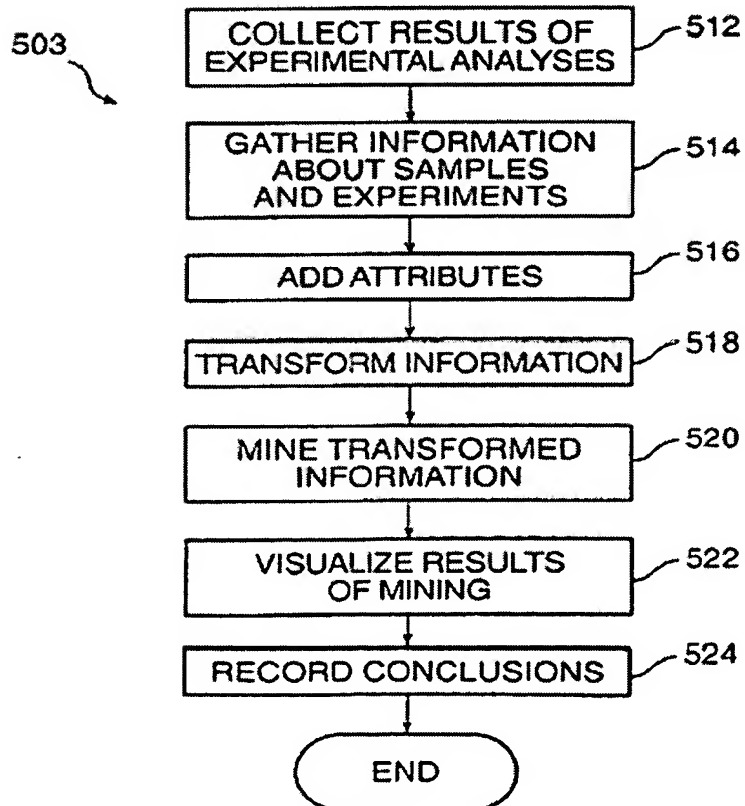


FIG. 5B

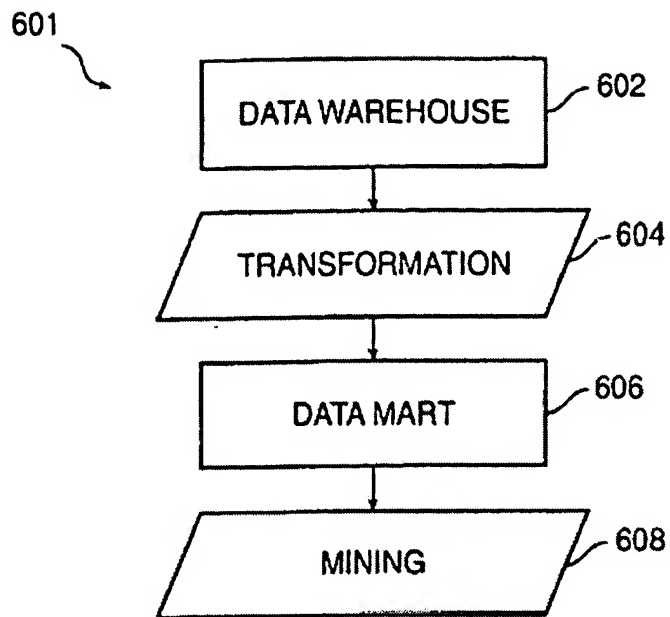


FIG. 6A

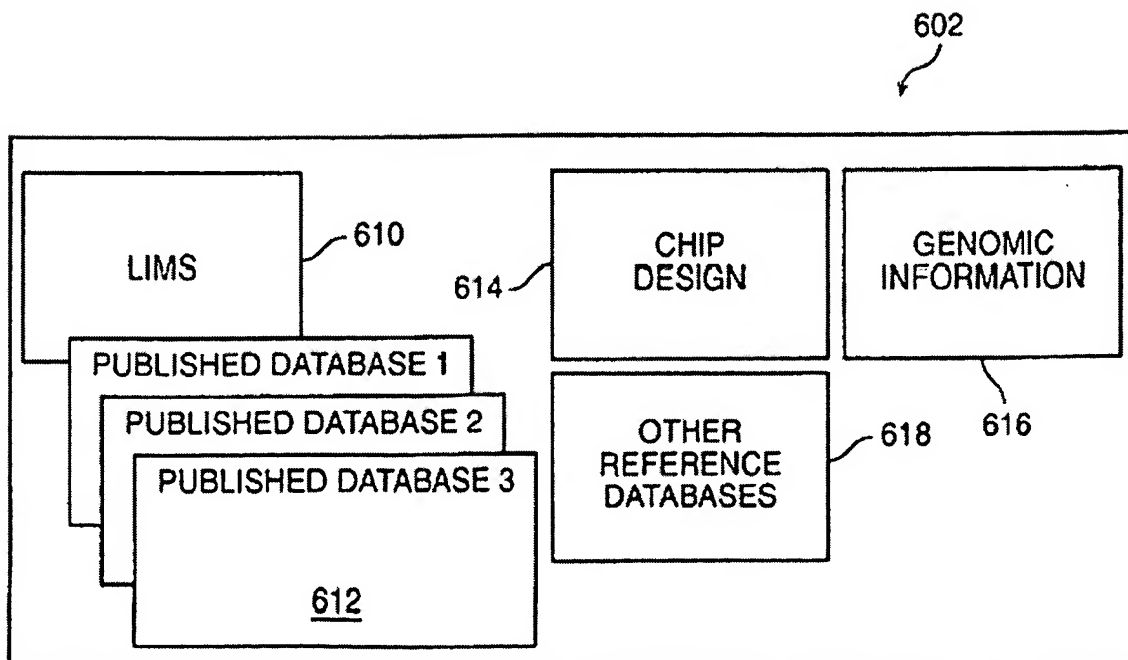


FIG. 6B

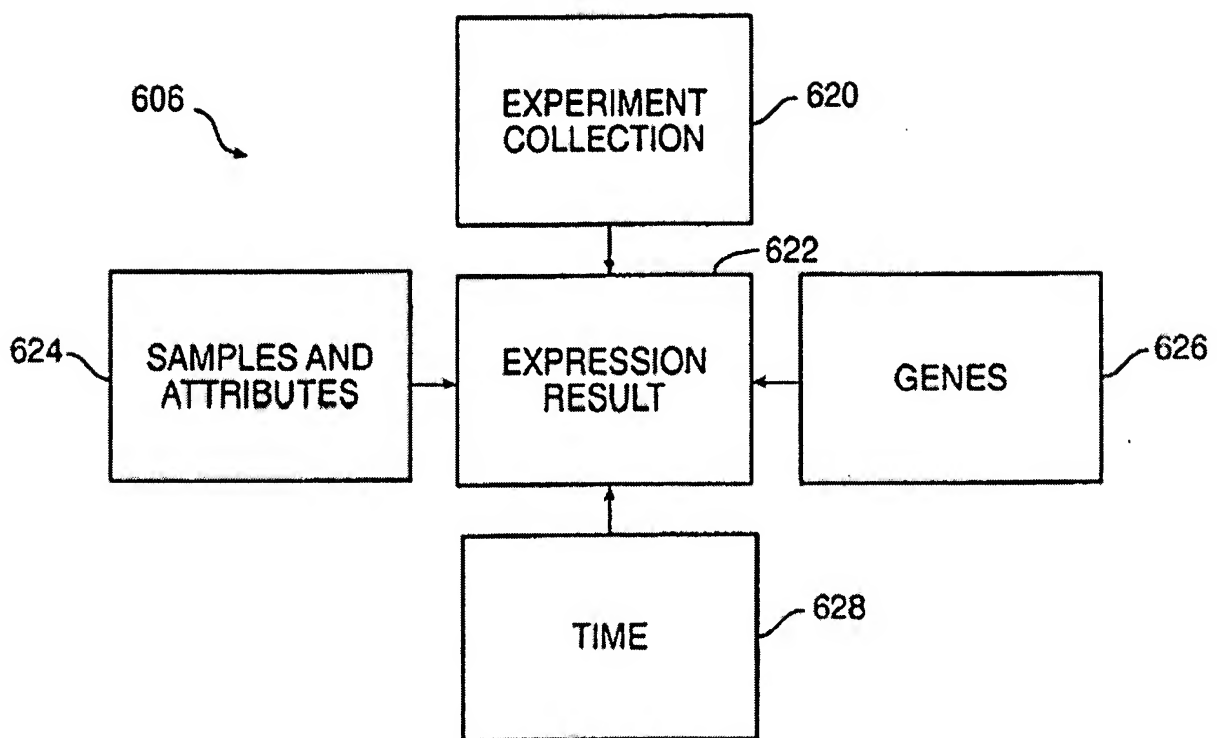


FIG. 6C

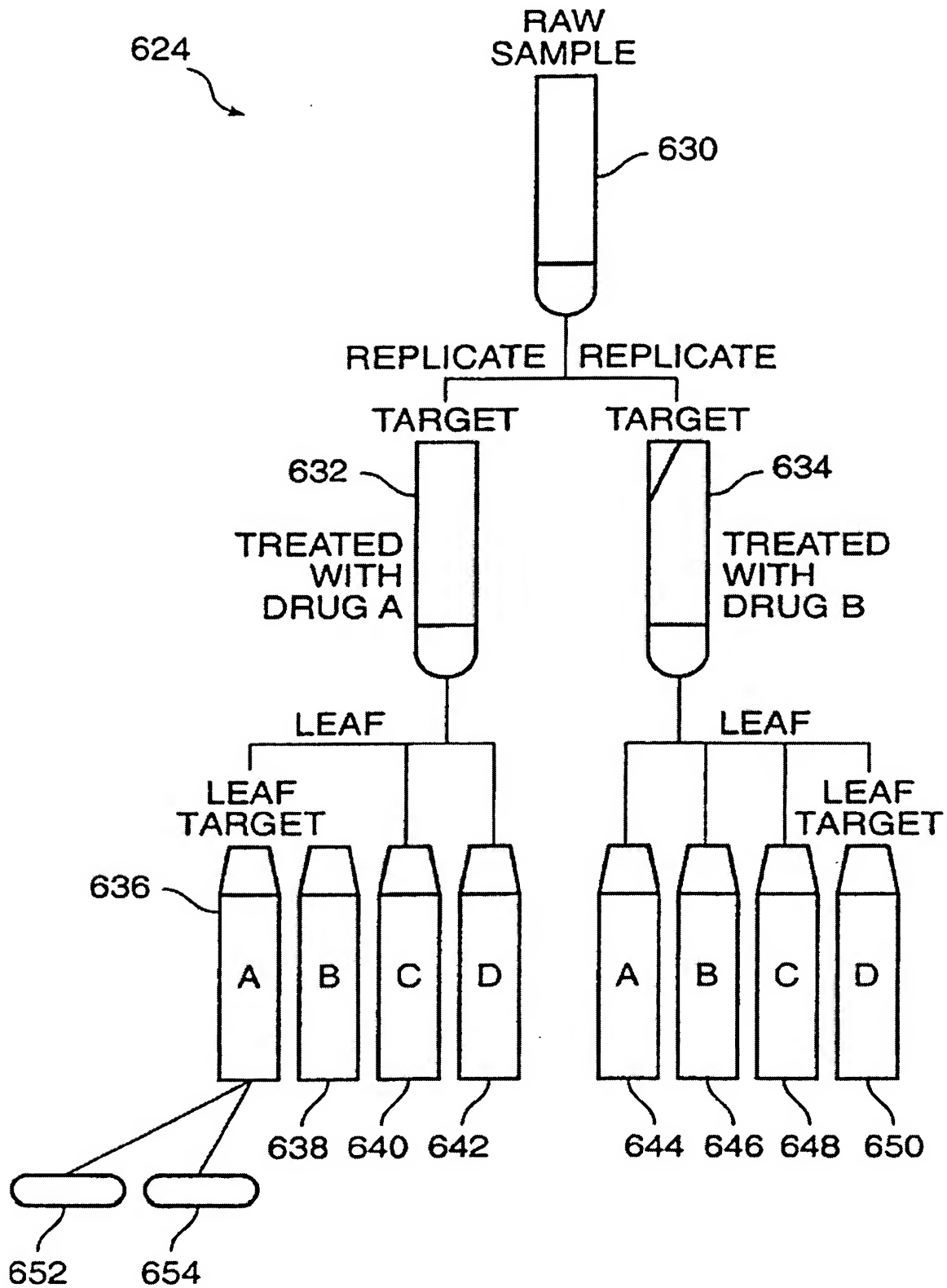


FIG. 6D

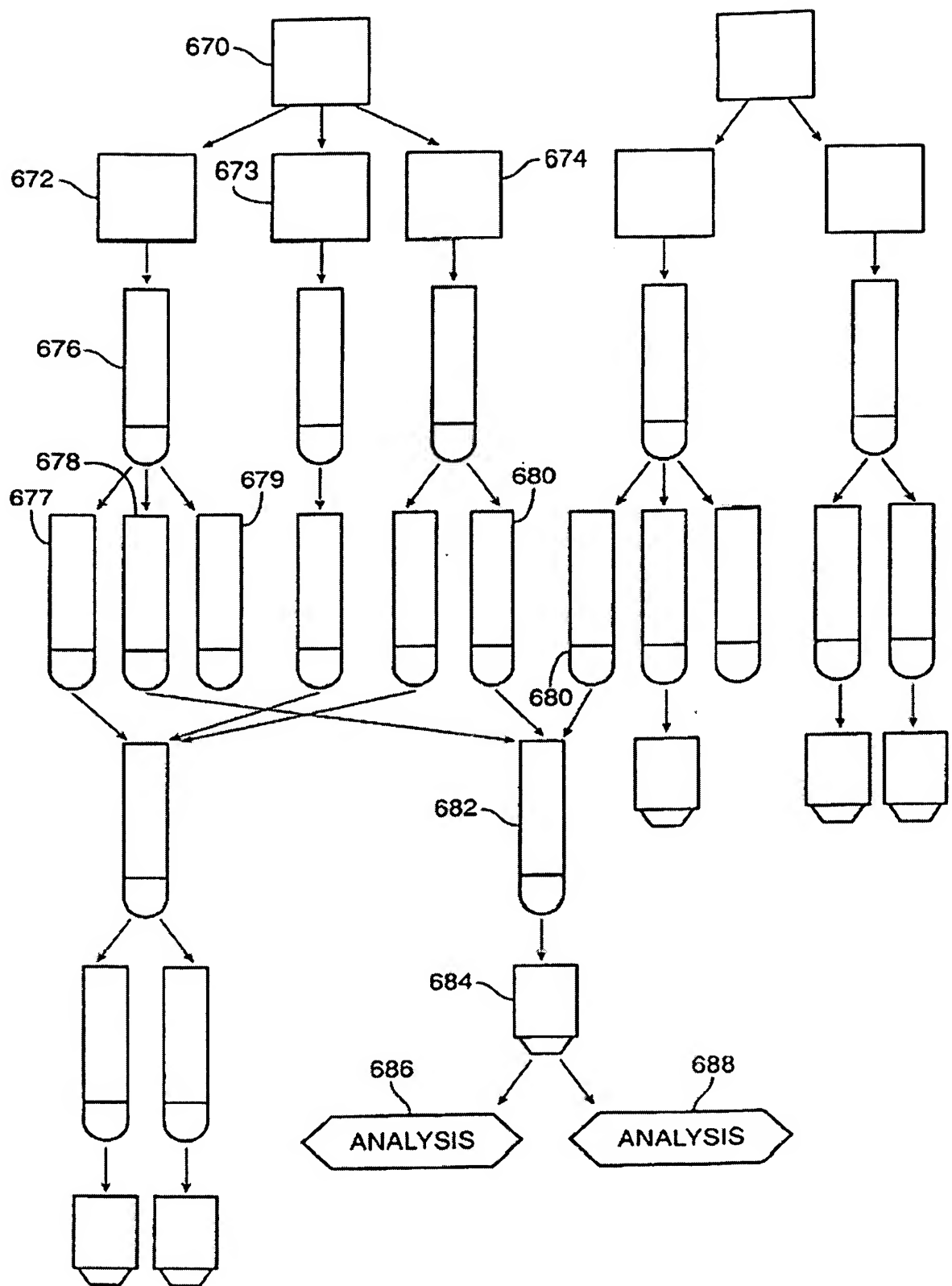


FIG. 6E

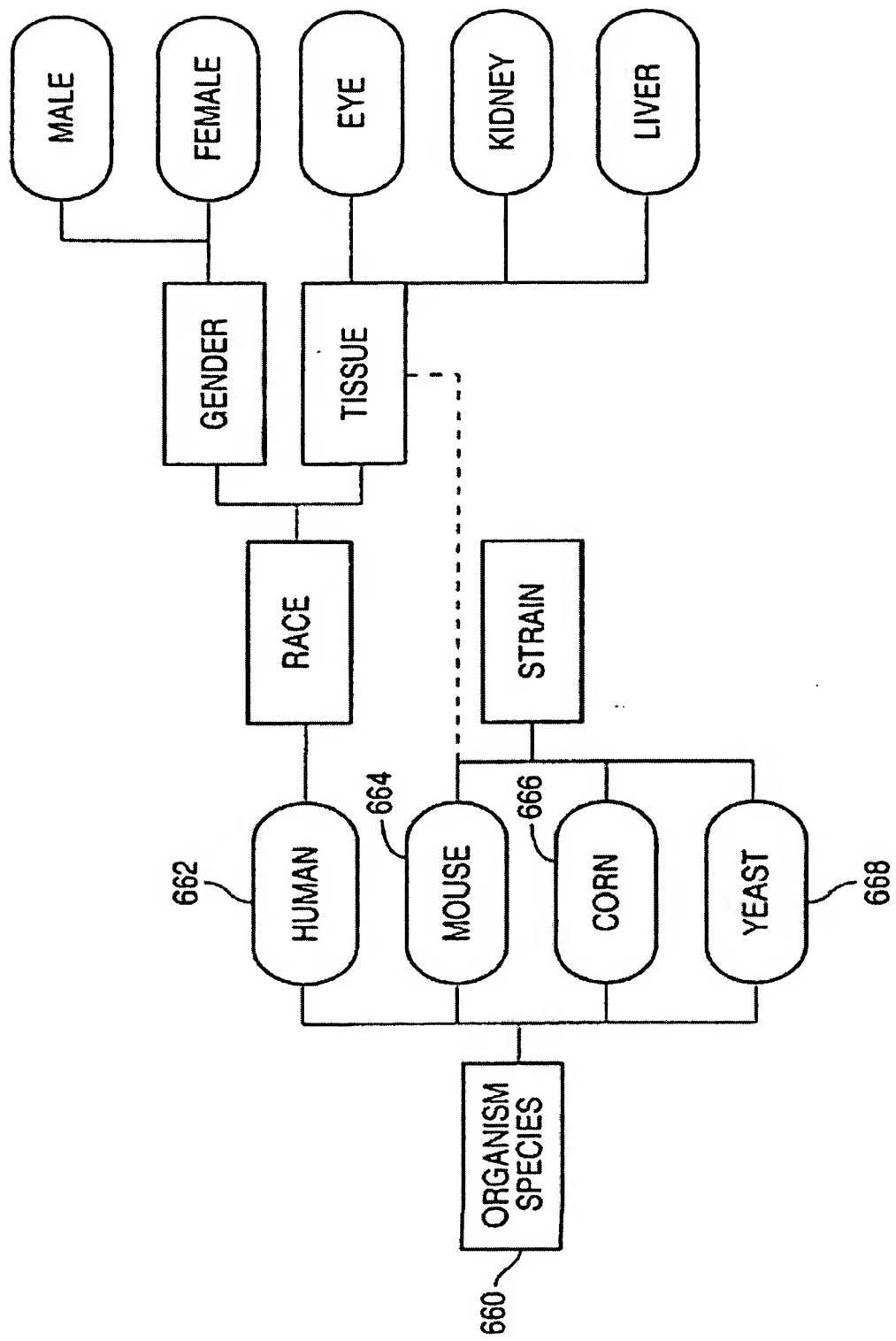


FIG. 6F

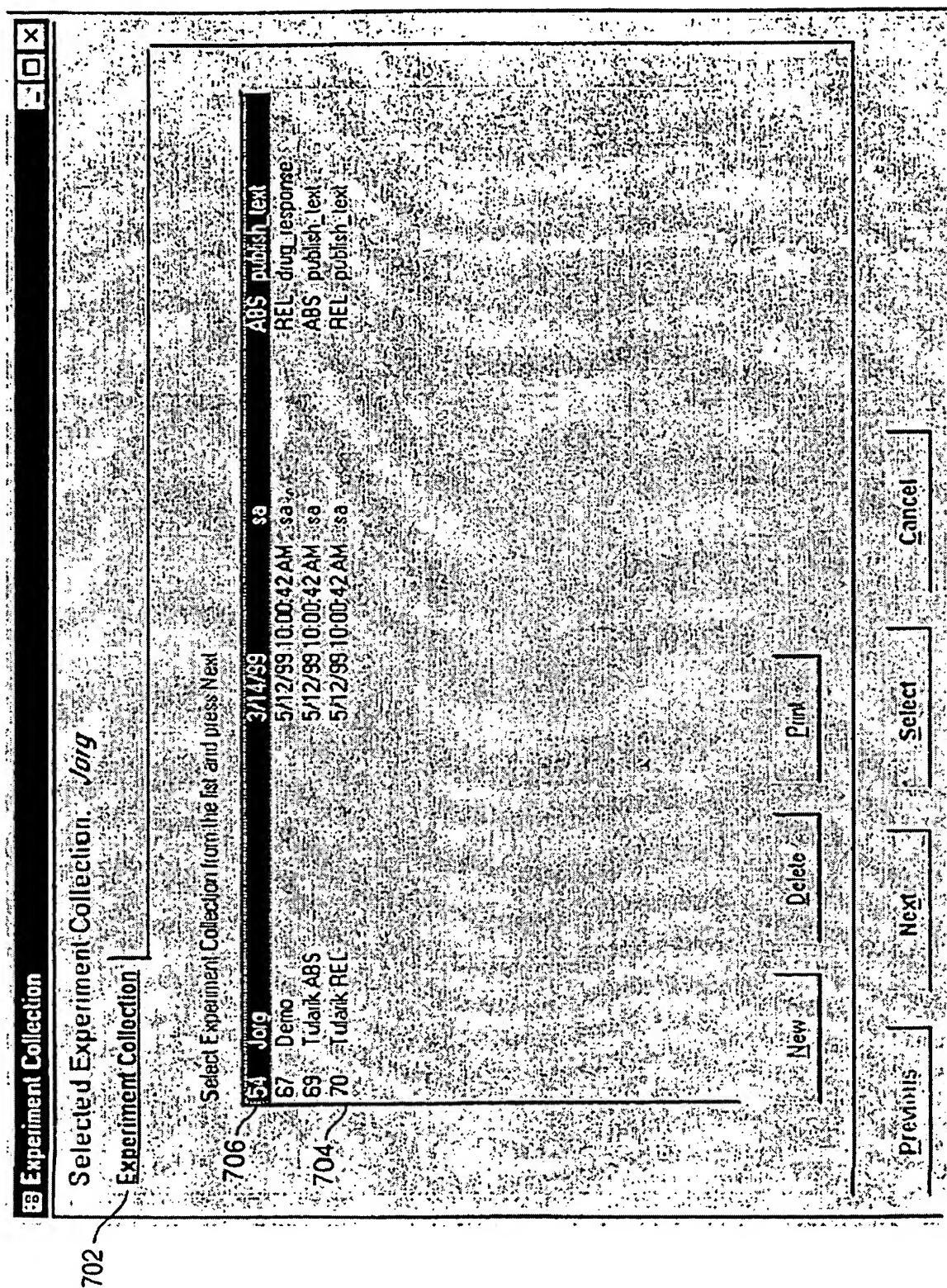


FIG. 7A

Experiment Collection

Selected Experiment Collection: Demo

Experiments

Experiments Selected

734

| expt | dat | file name |
|------|-----|--|
| 1 | | \\swdevalpha\GCLims\Data\980804D-3a.DA |
| 4 | | \\swdevalpha\GCLims\Data\9801268-3g.DA |
| 5 | | \\swdevalpha\GCLims\Data\9801268-2b.DA |
| 6 | | \\swdevalpha\GCLims\Data\9801268-2c.DA |
| 7 | | \\swdevalpha\GCLims\Data\9801268-2d.DA |
| 8 | | \\swdevalpha\GCLims\Data\9801268-2e.DA |

738a

738b

Experiments Available

736

| expt | dat | file name |
|------|-----|--|
| 9 | | \\swdevalpha\GCLims\Data\9801268-2f.DA |
| 10 | | \\swdevalpha\GCLims\Data\9801268-2g.DA |
| 11 | | \\swdevalpha\GCLims\Data\9801268-2h.DA |
| 12 | | \\swdevalpha\GCLims\Data\9801268-3a.DA |
| 13 | | \\swdevalpha\GCLims\Data\9801268-3d.DA |
| 14 | | \\swdevalpha\GCLims\Data\9801268-2a.DA |
| 15 | | \\swdevalpha\GCLims\Data\980827dE.DA |
| 16 | | \\swdevalpha\GCLims\Data\980826A2.DA |
| 17 | | \\swdevalpha\GCLims\Data\980826A3.DA |
| 18 | | \\swdevalpha\GCLims\Data\980826A1.DA |
| 19 | | \\swdevalpha\GCLims\Data\980826A4.DA |
| 20 | | \\swdevalpha\GCLims\Data\980826A6.DA |

744

Columns

746

Value

748

750

Filter

Reset

Previous

Next

Select

Cancel

FIG. 7B

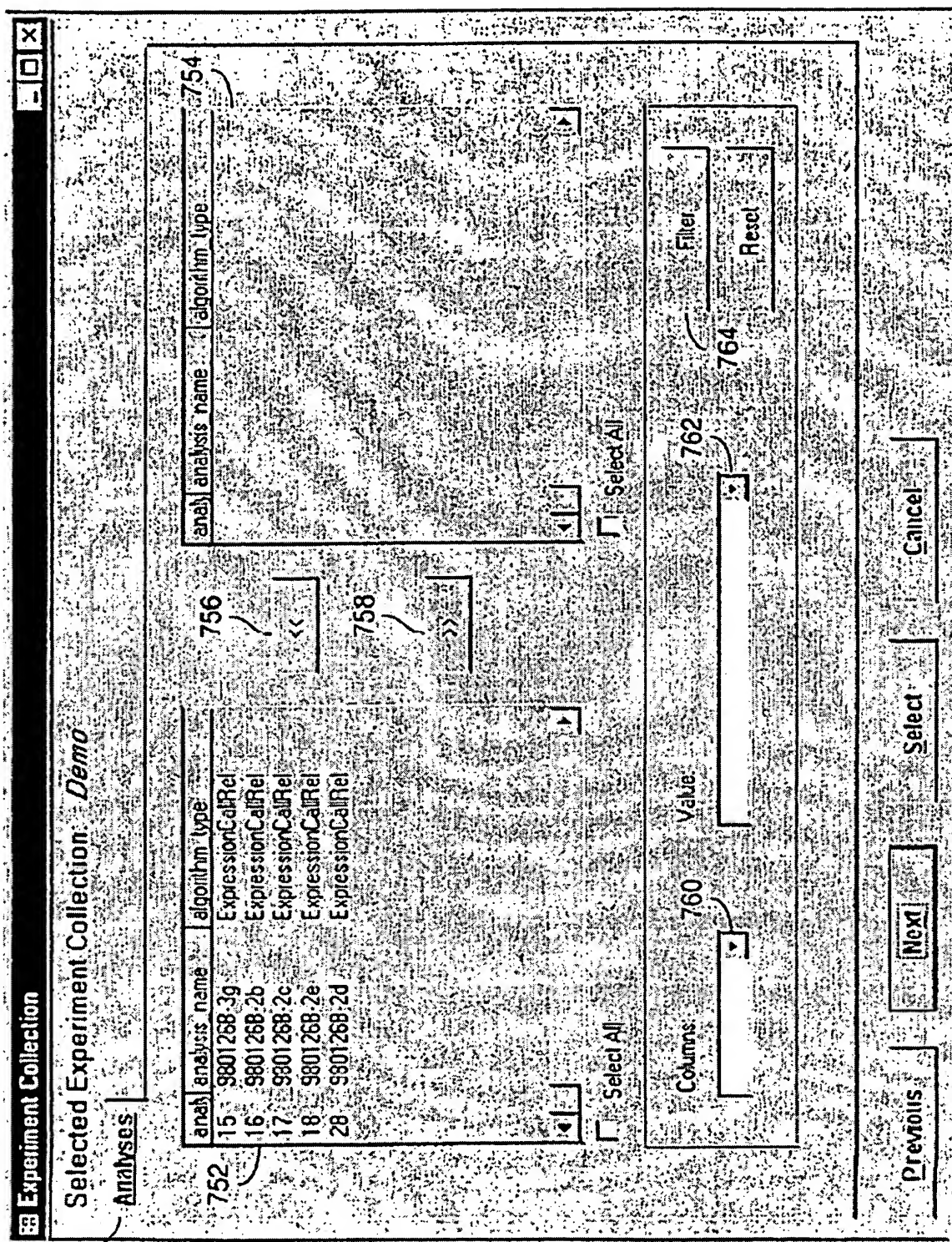


FIG. 7C

Experiment Collection

Selected Experiment Collection: Demo

Samples

Child Sample Item

| sample_item | type | derivation |
|------------------|------|------------|
| 7.8hr-control-ab | Leaf | True |

Sample Item Type

771

Parent Sample Item

| sample_item | type | derivation |
|-------------|-------------------|------------|
| 632 | 30min-control4-ab | |
| 633 | 30min-HIV-ab | |
| 634 | 2hr-control-ab | |
| 635 | 2hr-HIV-ab | |
| 636 | 4hr-hiv-ab | |
| 637 | 4hr-control-ab | |
| 638 | 8hr-control-ab | |
| 639 | 8hr-hiv-ab | |
| 640 | 16hr-control-ab | |
| 641 | 16hr-hiv2-ab | |
| 642 | 24hr-control-ab | |
| 643 | 24hr-hiv-ab | |
| 644 | 48hr-control2-ab | |
| 645 | 48hr-hiv-ab | |
| 646 | 72hr-control3-ab | |
| 647 | 72hr-hiv2-ab | |

Attribute

772

| attribute | type |
|-----------|-------------|
| 8 hr | Time Points |
| Project1 | Project |

774

Previous

775

Select

Next

Cancel

Manage Sample Attribution

Show in Visio

FIG. 7D

FIG. 7E

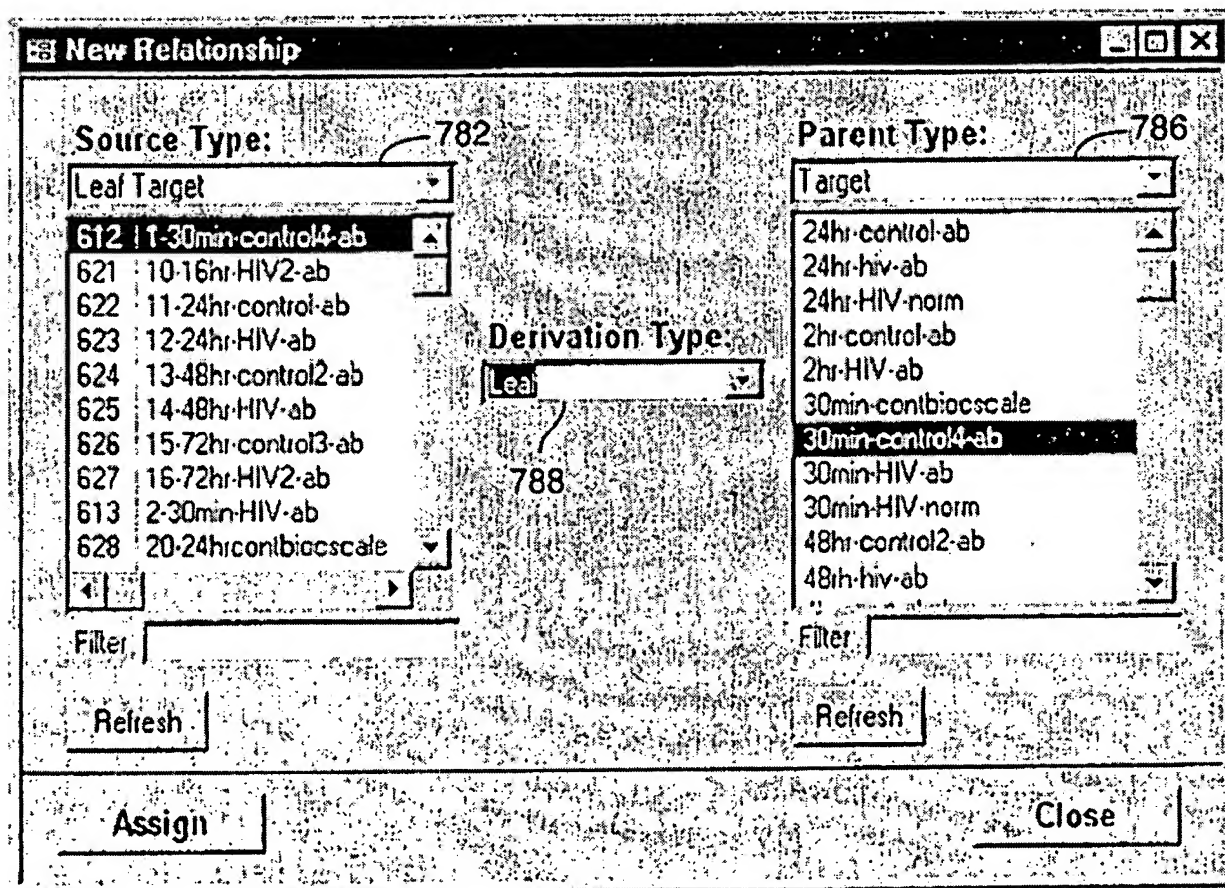
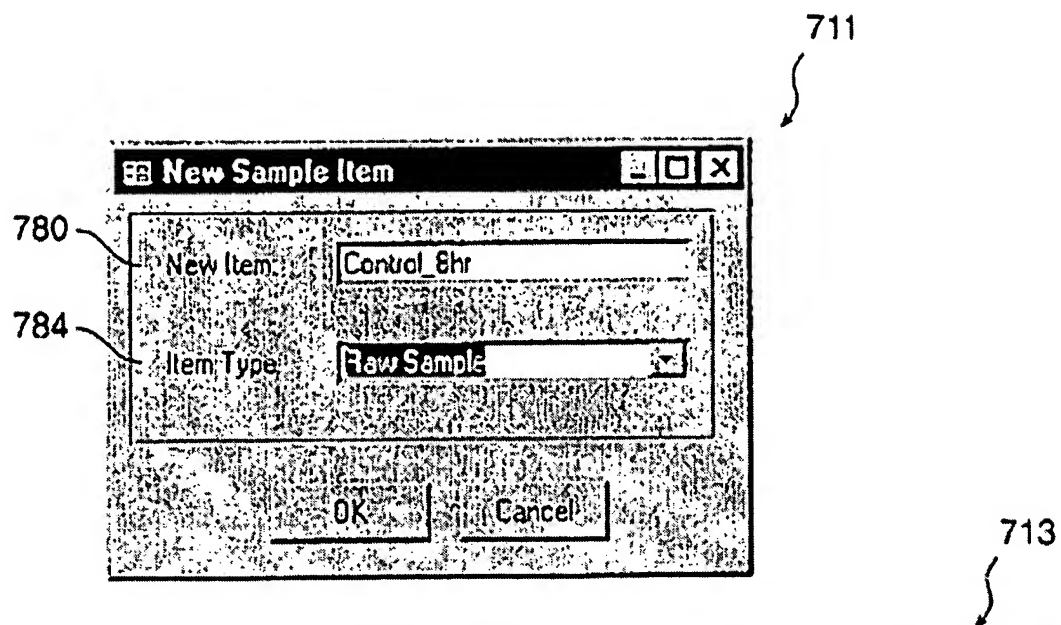


FIG. 7F

Experiment Mining

Select Group: 798

Project1

Saved Criteria

Save Criteria

| SAMPLE ITEM | BIO ITEM | NUMBER POS | CALL | NUMBER |
|-------------------|----------|------------|-------|--------|
| 30min-control4-ab | | | = 'p' | |
| 30min-HIV-ab | | | = 'p' | |

Record: 14

3

of 3

Gene Set: DEFAULT

Edit Gene Sets

Find all genes expressed in

100

% of the experiments in the group

801

Execute

Close

FIG. 7H

| Jag - Query ID: 1 | | | | | | | | | |
|-------------------|-----------------|---------------|------------------------|------|-----------------|--------------|------------|------------|--|
| Selected Data | | Scanner | Gene/Experiment Series | | Multiple Values | | | | |
| 804 | | 805 | 806 | 807 | 808 | 809 | | | |
| LEAF_PARENT | | EXPERIMENT | BIOTERM | CALL | AVG LOG RA | AVG DIFF INT | NUMBER POS | NUMBER NEG | |
| 30min-control4 | 1-30min-control | AB000449_at | P | | 376 | 8 | 1 | | |
| 30min-control4 | 1-30min-control | AB000460_at | P | | 984 | 12 | 2 | | |
| 30min-control4 | 1-30min-control | AB000468_at | P | | 1108 | 14 | 0 | | |
| 30min-control4 | 1-30min-control | AB001105_at | P | | 141 | 7 | 1 | | |
| 30min-control4 | 1-30min-control | AB001325_at | P | | 734 | 11 | 3 | | |
| 30min-control4 | 1-30min-control | AB002356_s_at | P | | 340 | 12 | 2 | | |
| 30min-control4 | 1-30min-control | AB002533_at | P | | 19048 | 19 | 0 | | |
| 30min-control4 | 1-30min-control | AB002559_at | P | | 1151 | 14 | 2 | | |
| 30min-control4 | 1-30min-control | AB003102_at | P | | 918 | 14 | 0 | | |
| 30min-control4 | 1-30min-control | AB003177_at | P | | 542 | 14 | 1 | | |
| 30min-control4 | 1-30min-control | AB003693_at | P | | 94 | 7 | 1 | | |
| 30min-control4 | 1-30min-control | AB004884_at | P | | 552 | 12 | 0 | | |
| 30min-control4 | 1-30min-control | AB006190_at | P | | 258 | 12 | 2 | | |
| 30min-control4 | 1-30min-control | AC000064_cds | P | | 174 | 10 | 2 | | |
| 30min-control4 | 1-30min-control | AC002045_xpt2 | P | | 1215 | 17 | 1 | | |
| 30min-control4 | 1-30min-control | AC002115_cds | P | | 6029 | 18 | 1 | | |
| 30min-control4 | 1-30min-control | AC002115_ma2 | P | | 557 | 13 | 2 | | |
| 30min-control4 | 1-30min-control | AD000092_cds | P | | 795 | 15 | 3 | | |
| 30min-control4 | 1-30min-control | AF000177_at | P | | 522 | 11 | 0 | | |
| 30min-control4 | 1-30min-control | AF006084_at | P | | 633 | 13 | 3 | | |
| 30min-control4 | 1-30min-control | AF006087_at | P | | 661 | 12 | 1 | | |
| 30min-control4 | 1-30min-control | AF007875_at | P | | 559 | 13 | 3 | | |
| 30min-control4 | 1-30min-control | AF009368_at | P | | 1256 | 13 | 2 | | |
| 30min-control4 | 1-30min-control | AF015913_at | P | | 2151 | 18 | 0 | | |

FIG. 71

| Selected Data | Scanner |
|-------------------------|---------|
| Select Value to display | |

Gene/Experiment Series

Multiple Values

Select Value to display:

Select Axes X points in the order you want to see them.

AVG DIFF INTENSITY:

30m-con-04-ab

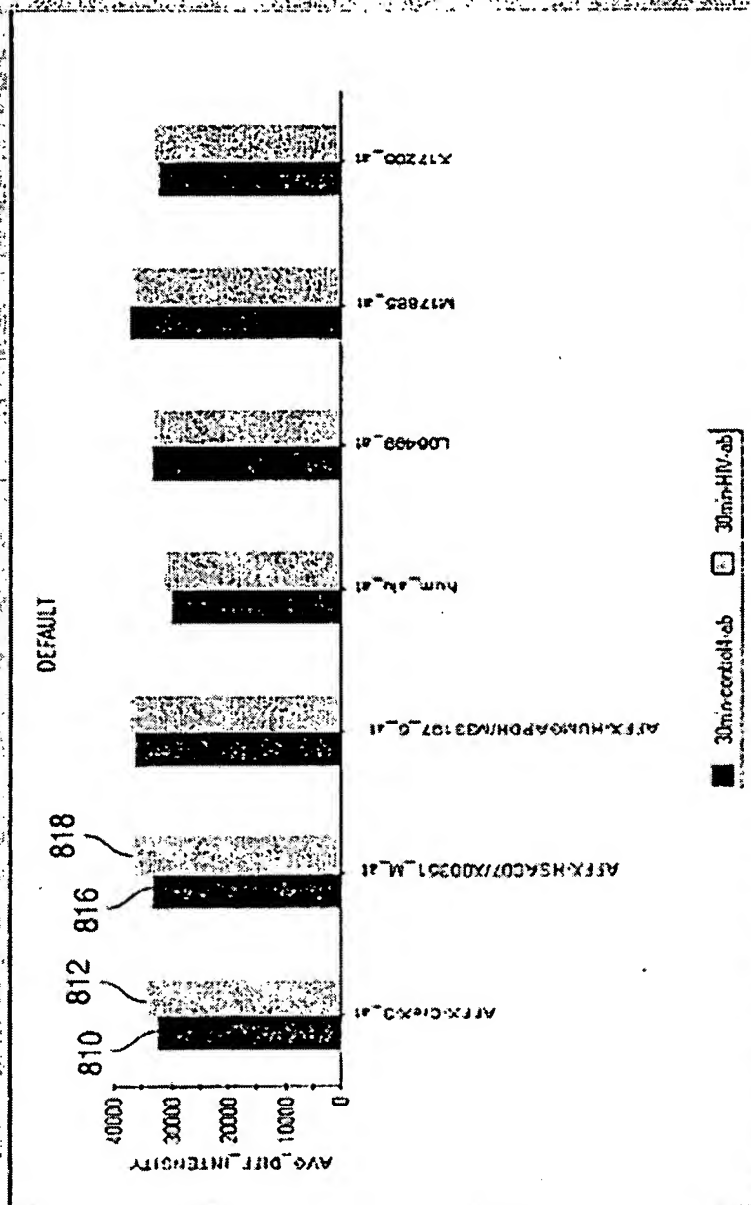
Can-con't

८३५

92

Save: chat

Save Chat



Prevlois

Next

Finish

General

FIG. 7J

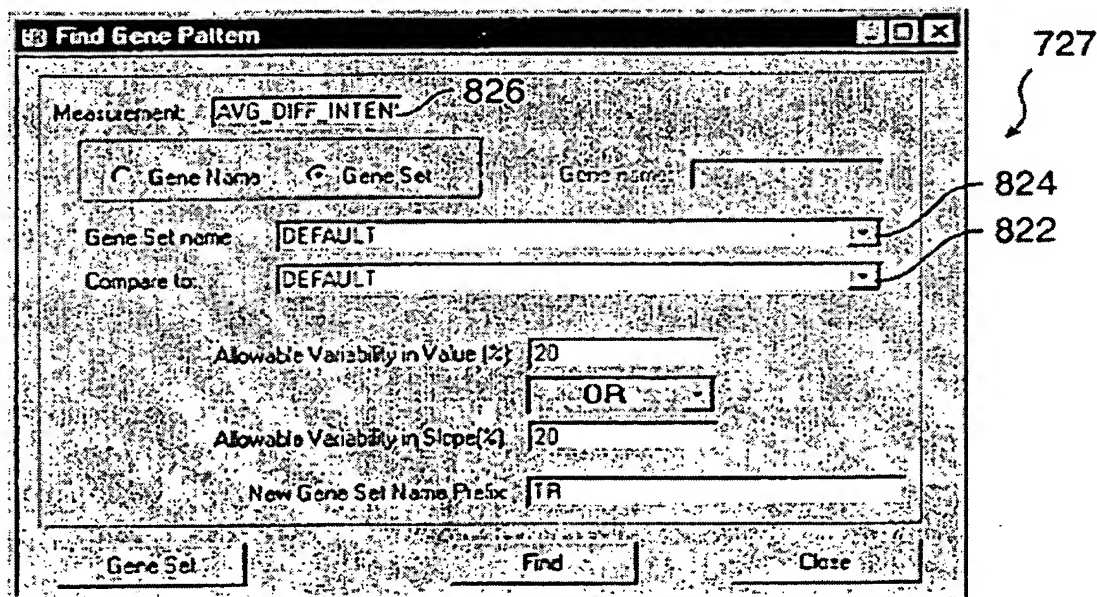
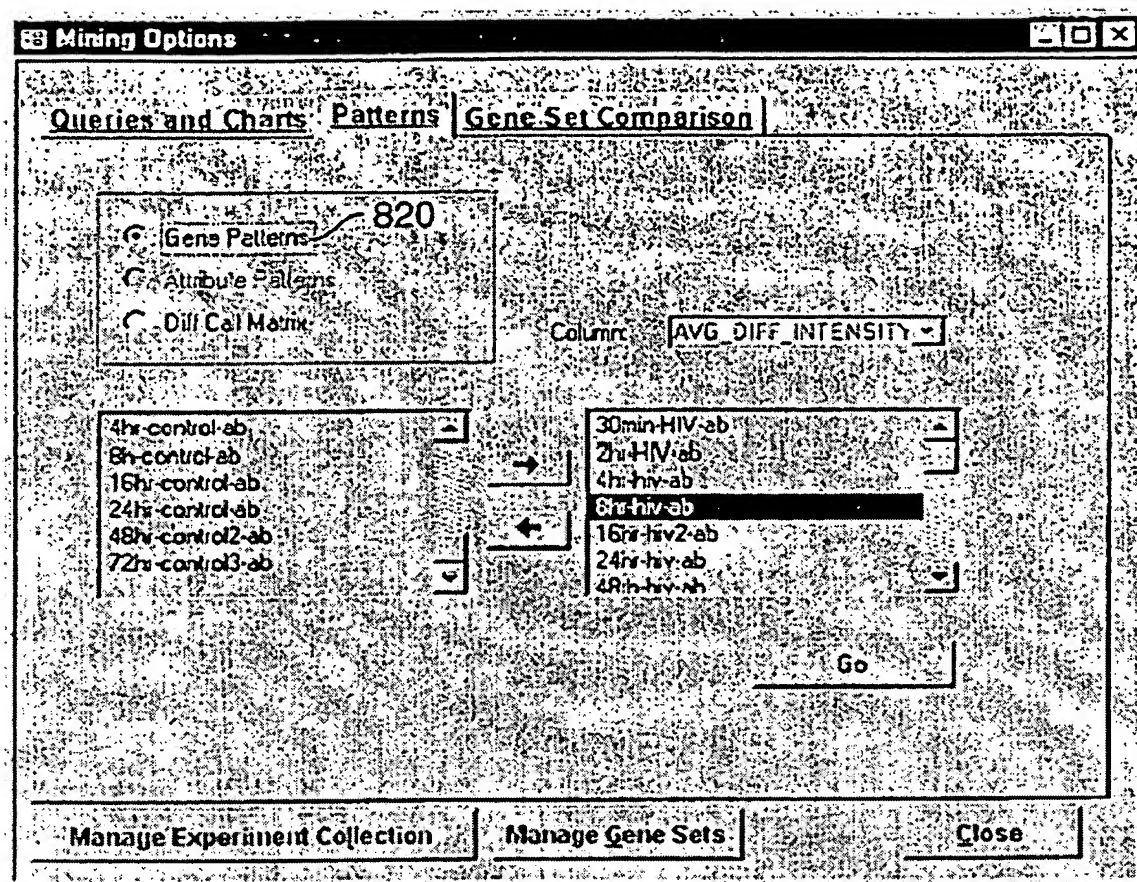


FIG. 7L

Mining Options

Queries and Charts | Patterns | **Gene Set Comparison**

Select Gene Sets to Compare:

| | | | |
|------|-----------------------------------|----|--------------------|
| 825 | DEFAULT | sa | 2/11/99 1:13:30 PM |
| 2793 | HepG2_IL_1-NC(1o6) | sa | 6/4/99 7:23:29 PM |
| 2794 | HepG2_IL_1a-NC&ID>(1o6) | sa | 6/4/99 7:30:45 PM |
| 2795 | HepG2_IL_1a-NC&ID<-100(1o6) | sa | 6/4/99 10:24:23 PM |
| 2796 | HepG2_IL_1ab-NC(1o9) | sa | 6/4/99 10:44:30 PM |
| 2797 | HepG2_IL_1ab-NC&ID+100(1o9) | sa | 6/4/99 11:25:47 PM |
| 2798 | HepG2_IL_1a-NC&ID-100(1o9) | sa | 6/4/99 11:27:58 PM |
| 2799 | HepG2_IL_1a-NC_ID&AD+100(1o9) | sa | 6/5/99 12:10:53 AM |
| 2800 | HepG2_IL_1a-NC&ID-100&AD+100(1o9) | sa | 6/5/99 12:29:33 AM |
| 2802 | Overlap | sa | 6/5/99 12:49:47 AM |
| 2803 | HepG2_IL_1ab-NC&ID100(1o9) | sa | 6/5/99 12:54:45 AM |
| 2804 | HepG2_IL_1a-DMD-100&FC-1(1o9) | sa | 6/5/99 1:32:16 AM |
| 2805 | HepG2_IL_1ab=ID+100&FC+1(1o9) | sa | 6/5/99 1:35:04 AM |
| 2806 | HepG2_IL_1a-CMD100&FC1(1o9) | sa | 6/5/99 1:39:43 AM |
| 2807 | HepG2_IL_1a-CMD100&FC1(1o9) | sa | 6/5/99 1:39:54 AM |
| 2808 | HepG2_IL_1ab=C&ID100&FC1(1o9) | sa | 6/5/99 1:43:19 AM |
| 2809 | HepG2_IL_1ab=C&ID100&FC1(1o9) | sa | 6/5/99 1:45:52 AM |
| 2810 | HepG2_IL_1ab=C&ID100&FC1(1o9) | sa | 6/5/99 1:50:31 AM |
| 2811 | HepG2_IL_1ab=C&ID100&FC1(1o9) | sa | 6/5/99 1:51:37 AM |

0 selected

Refresh Select

Manage Experiment Collection Manage Gene Sets Close

fmGeneSets_Compare_Form

| | HepG2_IL_1-NC(1o6) | HepG2_IL_1a-NC&ID>(1o6) | HepG2_IL_1a-NC&ID<-100(1o6) |
|-------------------------------------|-------------------------------------|-------------------------------------|-----------------------------|
| <input checked="" type="checkbox"/> | <input checked="" type="checkbox"/> | <input type="checkbox"/> | 836 |
| <input type="checkbox"/> | <input checked="" type="checkbox"/> | <input checked="" type="checkbox"/> | 838 |
| <input checked="" type="checkbox"/> | <input type="checkbox"/> | <input checked="" type="checkbox"/> | |
| <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | |

Record: 14 of 3

Enter name of new Gene Set Here: Geneset 2 of 3

Create Gene Set Close Form

FIG. 7M

Genes

Select Gene Set: **DEFAULT**

Double-click on
Accession to go
to Entrez

Copy My Gene Set Copy Other User Gene Set New Gene Set Delete Gene Set

ACCESSION: PROBE_PAIR_NAME DEFINITION:

| ACCESSION | PROBE_PAIR_NAME | DEFINITION |
|-----------|-----------------|---|
| A28102 | A28102_at | Human GABAA receptor alpha-3 subunit |
| AB000114 | AB000114_at | Human mRNA for osteomodulin, complete cds |
| AB000115 | AB000115_at | Human mRNA, complete cds |
| AB000220 | AB000220_at | Human mRNA for semaphorin E, complete cds |
| AB000381 | AB000381_s_at | Human DNA for GPI-anchored molecule-like protein, complete cds |
| AB000409 | AB000409_at | Human mRNA for MNK1, complete cds |
| AB000410 | AB000410_s_at | Human HOGG1 mRNA, complete cds |
| AB000449 | AB000449_at | Human mRNA for VRK1, complete cds |
| AB000450 | AB000450_at | Human mRNA for VRK2, complete cds |
| AB000460 | AB000460_at | Human mRNA, clone RES4-22B, complete cds |
| AB000462 | AB000462_at | Human mRNA for SH3 binding protein, clone RES4-23A, complete cr |

Add Genes Remove Genes Total Genes: **35620**

Record: 1 of 35620 (Filtered)

Close

FIG. 7N

Updated Gene Set

Accession:

Definition:

Filter

Reset

| | | |
|--------|----------------|---|
| Apr | 1-Apr | |
| M95678 | 1000M95678 | Homo sapiens phospholipase C-beta-2 mRNA, complete cds |
| T51617 | 1000T51617 | yb27h05.s1 Homo sapiens cDNA clone 72441 3' |
| M83353 | 1001M83353 | Human plasma membrane calcium-pump ngATPase (PMCA4) mRNA, complete cds |
| R06239 | 1001R06239 | Homo sapiens DNA-binding protein (PO-GA) mRNA, complete cds |
| L14922 | 1002L14922 | Human placental protein (PP1) mRNA, complete cds |
| M32402 | 1003M32402 | yb28d09.s1 Homo sapiens cDNA clone 72497 3' |
| T51944 | 1003T51944 | Human peroxisome proliferator activated receptor mRNA, complete cds |
| L02932 | 1004L02932 | |
| T51849 | 1004T51849_f | |
| T51849 | 1004T51849_i | |
| T51849 | 1004T51849_r_j | |
| L07592 | 1005L07592 | Human peroxisome proliferator activated receptor mRNA, complete cds |
| T51856 | 1005T51856 | yb54g11.s1 Homo sapiens cDNA clone 75044 3' similar to SP:GRPE_YEAST |
| T51955 | 1005T51955 | |
| U33635 | 1005U33635 | Human colon carcinoma kinase-4 (CCK4) mRNA, complete cds |
| T52003 | 1006T52003_i | |
| T52003 | 1006T52003_r | |
| T63591 | 1006T63591 | |
| T70046 | 1006T70046 | |
| R24080 | 1007R24080 | |
| T52014 | 1007T52014 | yb29q09.s1 Homo sapiens cDNA clone 72640 3' contains Alu repetitive element |

Remove Genes

Close

FIG. 70